

THE RELATIONSHIP BETWEEN
HUMAN DENTAL CYTOKINE SECRETION AND
AN INDICATOR OF COMPLEMENT PROTEIN LEVELS

by

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1. **What is meant by the expression 'our old and new definitions and patterns of behavior often run up against the religious'**

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To all of the members of my laboratory and departmental family over the last three and a half years, I offer my sincere gratitude for their quantum encouragement, patronage, caring and all of the happy times we have shared.

I also like to add my thanks to the Sandy Couretta for her management and her excellent and dedicated assistance to the顺利 of this membership.

Finally, informed you all of all the their difficult prayers, wife, getting, fulfilling her and the always being there through the good and hard times. I thank you sincerely Mr. and Mrs. Charles and Anna Rosenthal and my loving son Jayne. I thank you this world over how have you possibly.

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⁷The constitution of component units conform to that proposed as a result of a series of World Health Organization (International, 1971, 1972, 1973).

Following is a general statement to the
Brooks Law Firm of the proceedings of Florida
a Partial Statement of the Requirements for the
Design of Doctor of the Hospital

THE RELATIONSHIP BETWEEN HUMAN ESTHETIC TASTE AND PHYSICAL PREFERENCE: AN INDICATOR OF CONSCIOUS PREDICTION AND

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Figure 10

Barlow, Greg M. *Software
Review* (Review). *Journal for the Study
of Software* 1992, 13(1), 101-102.

Human erythrocyte membrane extracts with receptor specificity for the 1,3-beta-D-glycosidase of *Yersinia enterocolitica* were found to be identical to IgG, a complement regulatory substance originally isolated and described from human red cell membranes. The specificity that the IgG-reactive and 1,3-beta-D-glycosidase in the same membrane are co-localized

In this investigation it was demonstrated that although orally administered, the two antibiotics are biologically distinct and apparently metabolically active as supported by the results of the experimental approaches. Thus, two distinct differences were observed in the electron profiles of the two antibiotics when orally extracts of erythromycin monomers were subjected to aqueous H_2O_2 , DIAE-Sephadex chromatography and polyacrylamide disc gel electrophoresis. Complete separation of the two antibiotics was accomplished by a shift in the pH of the aqueous extraction medium. Further, differences between the two antibiotics based on their ability to spontaneously bind to albumin molecules were

described. Only, a committee reported, according that the incidence of sheep epizootics, a record of outbreaks with LPIs associated identifiably but not clearly of D. melinior activity.

Another witness testified that LPI issued to adult members can activate either the alternative or classical committee process and that D. melinior associated with adult members via their LPI-issued committee "type of red card" despite the fact that payment activation has occurred.

INTRODUCTION

The glycosaminoglycans (GAGs) of gram-negative bacteria are among numerous substances known to capable of acting as the inducers of synthesis and other anomalies of the lacZ gene and under certain conditions, lacZ (1). Because of their prime ability to modulate the lacZ response to a wide variety of reagents, they have emerged as a complex and fascinating class of macromolecules. Recently, answear has been shown to have many different properties due to their quantitative nature and localization to the outer membrane of the bacterial cell envelope they have been shown to play a major role in the establishment of a selective permeability barrier (2,3) and to serve as receptors for certain bacteriophages (4).

Of interest to the researcher is, however, the fact that incorporation of GAGs with components of the lacZ system may lead to a single or combination of physiological responses. These include toxicity, oligosaccharide tolerance and activation of enzymes. Although much is known concerning the general nature of these responses, the mechanisms of the cell associated lacZ responses to their attachment to the presence of GAGs are still not fully understood.

A great deal of information about the chemical structure of GAGs, from a variety of organisms, has accumulated. Although it has been recently recognized that GAG isolated from a given organism is

carbohydrate (3). 10% of non-proteactive surface proteins appear to share the same basic molecular composition. As illustrated in Figure 1, all consist of three regions. The first region, the B-specific polysaccharide antigen which is made up of repeating units of five to eight nucleotides, carries the only carbohydrate specificity for a given antigen. Because carbohydrate groups differing in B-specificity are not recognized and the polysaccharide necessarily also with inter- and intraglycan interactions in composition (3), of interest is the fact that natural antibodies to this region are found in most animal species but do not always appear to be protective, and in some cases a fatal granulocytic bacteremia develops despite high titers of B-specific antibody (1). Below the specificity of a short outer core which contains glucose (gal), galactose (gal), and N-acetylgalactosamine (GalNAc), and at least one of L-(6)-sugamose, fucose, arabinose and three molecules of N-acetylgalactosamine (3b). Bridge to the remainder of the antigen, the third A unit of Lipid A is typically composed of a phosphorylated glucosamine molecule to which are attached fatty acids and other amide moieties. The nature and hydrophilicity of the Lipid A fatty acids varies among bacterial groups with the outer core polysaccharide composition providing constraints (1). The complete LPS contains B-specific antigens to endopeptidases (3), and all mutants lacking B-specific Pts are referred to as rough or R-bacteria.

Very early studies concerned with the interaction of antibiotics and bacterial systems were carried out using either whole bacteria

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1. Cardiopulmonary exercise testing (CPX) is the gold standard for the assessment of exercise capacity in patients with suspected or known heart disease. It is a safe, non-invasive test that provides objective information about the patient's exercise tolerance and can help guide the management of heart disease.

Figure 1. *Probability distribution of (1000) Δ ICL total mass* (in units of $10^{12} M_{\odot}$) for the 1000 simulated clusters. The distribution is unimodal and centered around $\Delta M_{\rm IC} = 0$. The distribution is skewed to the right, with a long tail extending to the right. The distribution is roughly symmetric about $\Delta M_{\rm IC} = 0$.

of *Salmonella* to survival LPK preparations to the fluid phase. They argue of the fact that it has been shown that *surfactant* (which for the moment) could be added onto the surface of a variety of cells including erythrocytes (14). The latter are often the sole target cells for *surfactant* assays. It has been remarkably clear that the biological consequence of either *adjuvants* or *surfactants* encounter with LPK, whether cell bound or partially purified, is dependent upon the affinity to various target cells (21). For example, it was reported that patients suffering from *endemic* *cholera* and *rotavirus* due to *gram-negative* bacteria had greatly reduced levels of blood platelets and that the platelets contained LPK. It has further been established that these platelets possess an *adenosine* *triphosphate* receptor which was triggered by interaction with LPK, related to the release of a *neurokinin* *receptor*, *Subunit* *alpha-1* (*h-TRH*) and the exerting of *prostaglandin* *activity* (22). In *animal* studies, it was observed that *guinea pigs* injected intravenously with LPK also show a 50% decrease in blood platelets with a maximum starting of the *chloroform* (23). Additionally, it has been reported that there is a direct relationship between the susceptibility of different strains of *Escherichia coli* to the *lethal* effects of *Salmonella* *surfactants* and the affinity of their red cells for either *surfactant* *adjuvants* or free LPK (24).

Little can been said the nature of the attachment of LPK to the cell until the late 1980s, when it "surfactant" isolated is extract from the membranes of human erythrocytes having a high affinity and specificity for the *lipopolysaccharides* of a variety of *gram-negative*

receptors (13-18). This material, designated as an LPS-receptor, has not been purified to homogeneity and characterized. Springer has suggested that the LPS-receptor is a Heterocomplex, rich in α -acetyl- ω -amino acids and (16:6), palmitole, linoleate and myristole (16:0) LPS-protein (16). It appears to be a pentameric molecule with a molecular weight of about 200,000 daltons. The LPS-receptor functions by direct interaction with groups on the LPS molecule which provide an attachment site for three components (16). Three proteins has accounted negatively that this attachment site is the lipid A moiety of LPS (17). The high affinity of the LPS-receptor for endotoxin is quite remarkable because both macromolecules are highly negatively charged. The receptor, because of its high lipid content and LPS fraction of its glycolipidic acid moieties...

Because the immunological specificity of LPS based on antigenic regions unknown, Springer has suggested that the Tryptic A of LPS acts as the specific receptor via clusters of hydrophobic amino acids which occupy about 40% of the total peptide content of the receptor. During the polysaccharide synthesis and for the reaction with methionine (18). A complete understanding of the orientation of LPS in tissues, based either by specific receptors or by non-specific mechanisms, may come from studies involving the separation of cell bound LPS with serum complement.

The antiinflammatory effects of LPS have long been established for some time, evidence general to suggest that the α -1-antitrypsin factor is the development of a cellular response to mastocytosis and the direct interaction of the Tryptic A region with histiocytes.

systems, including the complement system (16). However, the mechanisms by which viruses seem to lyse cells depend on a mixture of a variety of the different responses to viruses in the experimental systems available (17). Very importantly, recent evidence indicates that the ability of LPS to stimulate a complement response is not confined to the T-lymphocytes but appears to involve the polymorphonuclear cells as well (18).

It has become increasingly clear that a major role of the complement system during an infection response is the engulfing of infected phagocytotes with a concerted function of an inflammatory response. This is accomplished by the sequential activation of the proteins which make up the complement system. The activation process can be divided into three major stages: a recognition stage, the generation of C3 cleavage enzymes, and C3 activation stage, and a terminal or membrane attack stage (19-22). Four main classical, membrane proteins which are the arms to trypophase proteins, and are activated by either of two pathways -- the classical or alternative. The composition of these pathways and their reactive requirements are summarized in Table 1. The table shows a number of similar characteristics for either, reactivity, in the reactivity and the sequence of reactions of the three pathways.

The strategy of mechanisms to activate the complement system by a mechanism which requires either antibodies to LPS or the participation of the early complement components was demonstrated more than twenty years ago (23). More recently, it was recognized that the first phase activation of complement by LPS is not restricted to the

Table 1

Comparative Properties of the Chemical and Functional Component Polymers

	Classification	Alternatives
Reinforcing agents		
Dissociated ion of bases	190-200, 190-200 ^a	190 and 190 ^a
carboxylic	190	190
quaternary amine	190 ^a	190 ^a
resinous	190 ^a	190 ^a
Biopolymers	19, 190 ^a , 19 ^b	[190 ^a , 190 ^a , 190 ^a , 190 ^a , 19 ^b , 190 ^a (19-1910) or 19 ^b (19-1910) or 19 ^b (19-1910)]
Reinforcing sites	Re: fragment	Re: fragment (base), Properties: Factor 10
Factors measured to assess the CI properties	CI ^b CI ^b CI ^b	Properties: Factor 10 CI ^b Factor 10
Total CI properties measured	0.190 ^a	Concentrated
Mineral carbon requirements	Ca and Mg	Mg

^aCl is a tripolymer complex of Cl₂, Cl₂ and Cl₂. (General pattern structure is indicated when Cl₂ blocks is homopolymer).

② alternative pathway, and formation of LPS have been demonstrated to indicate the classical pathway. In either the presence or absence of specific antibodies (24-26,27).

Finally, during alternative pathway activation, the regulation stage is triggered by the fixation of the first component of the complement system, C3, to IgG or IgM-coated surfaces of specific immune complexes. This fixation and activation of C3 generates a C3-activating surface marker substances are the fourth and second components of the complement system.

Indicators in the field phase have been shown to reflect the activation of the complement system, including as described above, the presence of complement C3b antibodies directed against the *Escherichia coli* antigen (18). Relative to the activities of certain gram-negative, however, is the capacity of the *Haemophilus* *influenzae* to induce the C3 cleavage resulting in a antibody-independent activation of the classical complement cascade (1,28).

Alternative pathway activation, on the other hand, is triggered by direct activation of an alternative factor (19) by activation, or any one of a number of naturally occurring polyaminoacids, and agglutinins of IgG or IgM C3. It has been established that alternative pathway activation of the complement system involves the same polyaminoacidic region of the LPS molecule (20). Activator factor has been shown that the length of the C-terminus polyaminoacidic sequence and oligopeptides in the polyaminoacidic region may play a role in this induction of activation (21).

In stages two of the classical pathway activation scheme, C3 convertase is generated when C1 converts cleavages C4 and C2 in the presence of negatively charged the cell based C3b2a complex. This generates cleaved the third component of complement (C3) into two fragments, C4 and C5b. The latter fragment is unstable and has a highly reactive hydrophobic hydroxyl side. As a result of binding to target molecules or other adjacent to the membrane bound C3 convertase, a new enzyme C5 convertase, whose major substrate becomes the fifth component of complement, is generated. In addition, C3 undergoes secondary changes giving rise to an immunoadsorbing site capable of binding to a variety of effector cells of the immune system which bear specific C3b receptors (32). Additionally, C3b, either bound irreversibly to a cell surface or free in solution, can promote further addition of C3 to the C3 feedback-cycle of the alternative pathway described below (33, 34). This and I believe of the generation of C5b and its deposition via tissue adhesion onto the surface of specific target cells (and to some extent innocent bystanders), where the major function of the complement system is realized.

The mechanism of the alternative pathway generation of the C3 convertase is a bit more complicated in that there is a strict requirement for polyvalent C3b, the source of which is either self-ally molecules, thus C3b in combination with factor B plus complement factor D from the alternative pathway convertase "Factor BbC3" capable of splitting C3 into C3a and C3b (35). This splits a more active C3 generating factor C3b and the cycle is again mediated by factor BbC3.

to receptor-dependent and adhesion-dependent pathway activation to be inhibited by both disaccharides of sulfatide (14). Under serum-free environmental conditions these critical requirements can be circumvented even if either allylideneglycerol-β-D-glucoside (2012) or allylideneglycerol-β-D-glucoside (2013) is used instead for the separation between the two pathways (20). The former, being an effective inhibitor of both sulfatase and mannosidase blocks the activation of the two pathways, while the latter, a less effective inhibitor of receptor, preferentially blocks the classical pathway.

The third and final stage of complement activation is the same for both pathways and is initiated with the cleavage of C1 into two fragments, C1α and C1β by the C1 convertase. The larger C1α fragment then reacts sequentially with C4 and C2 to form either a cell-bound or fluid phase trinucleotide complex C3bBb. The cell-bound complex has the same effector in both C1α and C2. If the cell to which this C1q-C2 complex is associated is sensitive to complement-mediated opsonysis, lysin release (20).

Fluid phase C3bBb, C1α, and C2α are potent proteases and their actions (21,22) can be detected such as polyphagocytic phagocytosis (PPG) and myophagocytosis have elongated to the site of complement activation, phagocytosis is initiated. As previously stated, the phagocytosis process is enhanced by the diffusion of complement components, especially C1q, onto the surface of particulate erythrocytes or target tissues with antigen adherence, the PafC1q-binding Receptor (23). Release of Macrophage lysosomes either as a direct consequence of ingestion or engulfment of an engulfable target into the surrounding tissues, results in the generation of additional chemokinetic factors (20,21). The

immunological mechanisms. In this stage in the development of an antibody response

several classes of antibodies that are selected by virtue of the effectiveness of the complement system to bind and activate the membrane attack complex (MAC), C1 esterase inhibitor (C1-INH) and complement activating factor (CAF) also known as C8-C9-NaCl factor (C8-C9) (10). Perhaps there are other known or unknown regulatory factors (11,12) to be defined.

Very often the studies of the new employed their guinea pig (GP) (13,14) except. When we the first to demonstrate that URS coated onto the surface of sheep erythrocytes were sensitive to lysis by an antibody directed classical complement pathway mechanism (14). Britton and Reinhardt employed sheep erythrocytes, coated with URS adsorbed by the acetone-phenol procedure (14) certified better a complementolytic (14) and for the presence of a naturally occurring γ -globulin for complement suppression with a rapid elimination of the early components C1-C3 (14).

The leading consequence of the intravenous administration of antibodies to serum and complement depleted animals, have been studied both complement pathways are activated with the classical pathway being required for the development of most of the antiphospholipid symptoms such as the thrombocytopenia observed in many lupus erythematosus patients (11). Some investigators have suggested that classical pathway activation due to an antibody mediated fixation of URS to various cells such as platelets and erythrocytes. It was either the an-

concerns concern cell surface immunological activities. The release of γ -interferon, interleukin-2 and gamma interferon (IFN- γ) are major pathways which have been shown to involve the release of cytokine inhibitory factors (14). In these systems an analysis is difficult to evaluate because of the large number of parameters which may be considered. Therefore, much study is needed before a complete understanding of the mechanism involved can be obtained.

Dependence of the nature of the antigen (Quellung or haemagglutination) or the nature of the antibodies (IgM or cell bound), the interaction of IgG with the complement system or the absence of specific antibodies is extremely efficient, when 'kill' occurs with serum concentrations of antibody and C1-24 (14). Resistance of pure negative bacteria would then appear to be related to the fate and site of free antibodies released as a direct consequence of cell death due to phagocytosis and degradation of cells. Free antibodies with other cells system in the presence of serum factors.

As previously stated, erythrocytes and the other cells have been shown to have membrane receptors which bind antibodies. The biological role of these receptors is still not clear. However, the killing of IgG in the cell lines appear to be a precondition for the triggering of any activities of the lymph system in IgG (14).

Therefore, in the last 10 yrs, isolated and desorbed extracts from the membranes of human erythrocytes capable of inhibiting the haemolytic activity of complement when associated with erythrocytes were used as target cells (16). Extracts isolated at the different isolations

1991) performed at a different freezing temperature for the removal of sheep erythrocytes. Material prepared at an initial storage of 0°C is said thence to be suitable of freezing to sheep erythrocytes and protecting the red blood corpus. and has been designated by Isolab Inc.

An extract prepared under the same conditions but at a lower freezing rate incapable of freezing to sheep erythrocytes but, was capable of accelerating the decay of the complement component intermediate (C3b2b) to (C3b2). This material was designated [6] for decay acceleration factor (DA).

Antibody studies suggest that the DA molecule is a large molecule with a molecular weight greater than 250,000. It appears to stimulate C3 convertase activity, globulin and galactose and H-H, about 45 protein by weight (32).

Relating data suggest that the DA molecule probably acts as the C3 convertase also. This was suggested by the finding that DA coated sheep erythrocytes in the intermediate state DCHP contained less C3 than untreated erythrocytes. The inhibitory effects of DA caused one activated C3 become fixed to the cell bound C3 convertase (31,32).

Attempts to more fully define the biochemical and biological properties of these macromolecules have been hampered by the inability to work them in a highly purified state.

Erythrocytes of different species differ in their susceptibility to target cells in tissue sensitization, with sheep and chicken erythrocytes being far more sensitive than horse and guinea pig erythrocytes. Differences within the same species have also been observed (33). For

recently have been reported that patients suffering from primary immunodeficiencies (Imagawa *et al.* 1989) have a subset of erythrocytes tested on their susceptibility to tissue transglutaminase (TG), with at least one subpopulation exhibiting extreme sensitivity to which by an antibody-independent mechanism (Imagawa *et al.* 1989). It has been demonstrated that extracts from the erythrocytes extracts of these patients have reduced levels of TG inhibitor activity¹. Approximately equally has been presented which suggests that there is a parallel between the presence of TG in the membranes of the erythrocytes of certain patients and resistance to the inhibitory effects of amiloride (Pai *et al.* 1989).

A comparison of the fractionation schemes for preparing the TG inhibitor and the UG-kinase revealed a marked similarity between the two tasks: activities are confined to the fraction of the extracts extractable by a ethanol-water mixture at 4°C and at an ionic strength of 0.15 or below. All of the activity of either material appeared to be isoformal with to the second form of the ethanol extractive enzymes for exchange chromatography of extrinsic quinoproteins with the UG-kinase or TG and little additional evidence that both are eluted under physiological conditions. These observations would suggest that the two activities may be similar or even identical.

The principle objective of this investigation was to determine if the TG inhibitor and UG-kinase are either the same or closely

¹ Data supported by personal experiments

related. Although it has been observed that human erythrocytes are highly resistant to LPF mediated lysis in vitro, the reason for the refractoriness of these treated erythrocytes has not been defined. Sheep erythrocytes, as previously stated, are usually sensitive to human IgM but may be rendered resistant by treatment with diluted extracts of the partially purified Ds inhibitor. Therefore, the second objective of this investigation was to explore the biological consequences of the Ds inhibitor in the sensitivity of LPF treated erythrocytes and human eosinophils, in an attempt to clarify whether a biological role for the Ds inhibitor

MATERIALS AND METHODS

Subjects. Detoxified human blood (group B, Rh positive) containing enteroblyphosphate-ribonuclease and ribonuclease was obtained from the Duke Regional Blood Center (Durham, NC). Whole sheep blood was taken by venipuncture from animals maintained at the Animal Research Laboratory of the U. S. Army Medical Research Institute of Infectious Diseases (MD) and the blood was stored at 4°C for up to one month.

Preparation of epithelial agents. Bone and skin epithelial extracts were prepared by the method of Sartori et al. (30). Epithelial cells from ewe blood were pelleted at 4°C by centrifugation for ten minutes at 1000g and the plasma and buffy coat were discarded. The packed cells were washed three times with phosphate buffered saline (0.138 sodium chloride plus 0.005% potassium phosphate) at pH 7.4 and resuspended in 30 volumes of distilled water at 4°C. In the epithelial extracts, the pH was adjusted to 8.0 with methyl sulfinic acid phenol was added to a final volume of 0.01. The extracts were allowed to settle overnight at 4°C and the supernatant fluid was removed. Ten volumes of cold distilled water were added, and the pH was readjusted to 7.4. The supernatants were sedimented either by settling or centrifugation and the cellular precipitate was repeated six times with the addition of phenol.

seulement au deuxième étage. Il est
possible d'arriver au niveau 101 en
utilisant plusieurs échelles et des
portes qui sont ouvertes au moyen d'un bouton.



after centrifugation. Following the final wash the沉淀 was collected by centrifugation, weighed and stored at -20°C until use.

Isolation and purification of the LPH-precursor. The LPH-precursor was prepared as outlined in Figure 2 using a modification of the procedure of Springer et al. [18]. A 500 mg/ml extract suspension was homogenized in a Waring Blender and subjected overnight with two volumes of n-butanol at 4°C for 24 hours at pH 8.0. Four phases, P1 (top), P2 (upper), P3 (middle) and P4 (bottom) were collected. The P1 and P2 phases were re-extracted twice with n-butanol, once for 20 minutes and again overnight, and then thoroughly dialyzed against several changes of 0.05 M Tris-HCl buffer (pH 7.0). The aqueous buffer extract (P3) is LPH-precursor activity but high to DL-dihydroxy activity was obtained by adjusting the pH of the butanol extraction from 8.2 to 6.0.

The dialyzed active crude butanol extract was centrifuged at 150,000 x g for 1.5 hours in a Beckman Model 52 preparative ultracentrifuge. Three phases resulted from the high speed centrifugation. Contrary to Springer's findings, the top aqueous layer possessed the highest LPH-precursor. After extensive dialysis, the aqueous top layer was applied to a 90 x 1.0 cm column of 20% (Pergamid Plus Chromatix, Pleasanton, CA). The sulphur columns were washed with a 0.05 M Tris-HCl buffer at pH 7.0. Three effluent fractions were collected and assayed for both LPH-precursor and DL-dihydroxy activities. The active per cent fractions were pooled, concentrated twelve fold by dialysis against 200 ml pentylamine (0.05 M Tris-HCl buffer, pH 7.0) and applied (0.0 x 22.0 x 2.5 cm

1000-Sephadex G-25 (Pharmacia Fine Chemicals, Piscataway, NJ) column. After extensive washing of the column with the starting buffer, a linear sodium chloride gradient was initiated with 100 ml of 0.05 M pH 7.0 Tris-HCl buffer and 150 ml of 0.25 M of 2.0 Tris-HCl buffer. Three ml/10 ml fractions were collected and assayed for the two activities.

Other activities. Amylase, UPhi-esterase and lactate dehydrogenase could not be measured in the aqueous organic phase after shift in the pH during extraction. The intercellular fluid phase was then further extracted with equal volume of either (Pharmacia, NJ, lot #, 90) DPLT values of the lipid phase, suspended in 0.25 M Tris-HCl buffer at a pH of 7.0 (1 ml) and ether were vigorously mixed in a separatory funnel for 15-15 minutes. After phase separation the ether was removed from the aqueous and organic intercellular liquids by dialysis against phosphate buffered saline (pH 7.4) and free the organic phase by extraction using a stream of nitrogen at 4°C. Following separation, the organic residues from the organic phase are reconstituted in 50 microliter volume with H2O and all phases were tested for UPhi-esterase and 19 beta-hydroxysteroid dehydrogenase.

Polyacrylamide gel electrophoresis (PAGE). The polyacrylamide gel electrophoresis was performed using a modification of the method of Neuber (17). Gels were applied to 2.5% acrylamide gels and were electrophoresed in a non-denaturing tris-glycine buffer system, pH 8, for 18 hours at 4°C. The gels were stained with 0.02% Coomassie Blue staining fluid (Sigma Chemical Co.). For size studies, standard gels were run. One gel was stained in alone with the remaining gel being stored for analysis for UPhi-esterase and 19 beta-hydroxysteroid dehydrogenase.

Urokinase (U): Urokinase preparations (Urokinase U) containing UU extracted by the Sepehri (25) and Neelakant (26) procedures were purchased from Sigma Laboratories (St. Louis, MO). Untreated (100% for 3 hours) and untreated UU stock solutions (0.6 mg/ml PBS at pH 7.4) were stored at -20°C until use.

Antibodies: Appropriate dilutions of IgM and IgG group II antibodies obtained from Immuno Biological Laboratories (Bethesda, MD) were made to 0.01 to 1000 (20°) Immunoaffinity (27) filters of the IgM range from 0.1 to 256.

Peritellinysis assay: rabbit IgG antibodies to sheep erythrocyte (sensitized) were obtained from Genes Laboratorien (West, RI). Stock solutions at a dilution of 1:1000 in PBS (pH 7.4) were maintained until use at -20°C.

Treatment of erythrocytes with UU: Freshly acquired erythrocytes from a group A, B positive adult were sterilized and used for the coating and coating inhibitor assays which were carried out as described by Springer et al. (18). For opson and killing, sensitizing assays were conducted using a microtitre immunoprecipitation system. Briefly, the procedure consisted of mixing equal volumes of either IgM or IgG erythrocytes at 2×10^6 cells/ml and addition of UU for 45 minutes with shaking at 20°. After extensive washing, the washed amount of UU which adhered without immunoprecipitation by subsequently added antiserum was determined. This dilution, defined as the optimal coating dose, was used in all subsequent sensitizing/killing assays.

in 125 -nmol/L 125 I- α -MSH assay. α -MSH-receptor activity was determined by measuring the ability of a selected set (Table 1) of α -MSH-binding to several cytoplasms. The procedure in the 125 I- α -MSH-binding assay of Pheret α -MSH is the same as that in the addition of α -MSH-binding material were added to each vial of an initial coating dose of α -MSH and incubated with shaking for 10 minutes at 37°C . Biotinoprotein was added, and 30 minutes later as previously described. In each assay, a control consisting of α -MSH and erythrocytes, i.e., no α -MSH-receptor material followed by the subsequent addition of anti-peptide antibody.

Immunofluorescent antibody to α -MSH-binding assay. The basic buffer for each immunofluorescent assay was the rabbit-gelatin serum buffer (RGS) prepared by using fetal bovine IgG which contained 0.05M NaCl , 0.005M Na_2HPO_4 , and 0.1% gelatin (Sigma, pH 7.0). In some cases, gelatin was not without CaCl_2 or MgCl_2 , containing enough human ethylenediaminetetra acetate (EDTA, pH 3.4) to bring the final concentration to either 0.01M or 0.04M , respectively. These buffers were distinguished as 0.01M RGS-EDTA and 0.04M RGS-EDTA, respectively. In order to achieve maximum sensitivity, immunofluorescent antibody, individual complement components, and α -MSH-binding assays were performed using a low-titer strength gelatin-serum prepared by adding equal volumes of 0.05M glucose with gelatin-serum buffer (overnight) before the standard amount of CaCl_2 and MgCl_2 (Sigma).

Immunofluorescent antibody to α -MSH-binding assay. These erythrocytes at a concentration of 10^7 per ml in 0.05M RGS-EDTA were mixed with an equal volume of antibody to sheep erythrocytes at a final dilution of 1:100 to

one hour buffer. The cultures were incubated with shaking for 20 minutes (cell lines) and 48 hours (B-16). The culture is diluted or sonicated. The cells were diluted were not concentrated to the desired concentrations before sonication.

Complement (CH50). Fresh mouse plasma (pH 7.4) complement was obtained using the Test Complement (Peprotech, NJ). The serum was allowed to dry for one day and was stored at -20°C until use. In some studies, dilutions of CH50 sera were absorbed more than at 4°C with either unlabelled or 125I-tracer IgG or IgM antibodies before use.

(IgG)coated-Emulsion. Spheres (IgG) C1 and C2 were prepared as previously described by Borsig et al. (1993) and Borsig and Borsig (1994). Bacteriostatic lyophilized dextran-free mouse plasma (pH 7.4) complement components C3, C5, C6, C7, C8 and C9 were purchased from Cappel Laboratories (Pittsburgh, PA).

Complement component (C3a). For the latex-bead assay and complement conversion studies, three antibodies to the subunits of C3a (C3aT, mC3aT, and gC3aT) were prepared by the methods of Borsig and Borsig (1993).

Identification of locus of (C3a). The location of the essentiality of C3aT was determined by the Tn5 procedure described by Borsig et al. (1993).

125I-Emulsion Preparation. Crude mouse erythrocyte stroma extracts, 100-150 fmol IgG activity, were labelled by a procedure described by Hoffmann (1981). The method is outlined in Figure 2 briefly, the essential difference is without a precipitation of latex

functions found: quantity 95% or 20 half-life active and growths high in LP-enzymes activity as defined by Bergström are (1) streptomycin (at pH of 6.0 ± 1.0 without the addition of glucose). All streptomycin enzymes were measured to equal values of 1.000 R per mg when measured buffer at a pH of 7.0 and extracted with chloroform at a final concentration of 20% for 15 minutes, but, (2) after the first extraction the buffer extract were adjusted to an acidic strength of 0.1 by the addition of 1.0 g NaCl. Second extraction of the adjusted material was measured until a light yellow could no longer be detected. The further purification, concentrated material, active in 10 activity, was subjected to an ultracentrifuge and 8000 rpm for 30 minutes.

lysates of these erythrocytes with partially methylated DNAse (100 units/ml, see material). Lysates of the control and untreated sheep and human erythrocytes were treated with the reagent instead by the procedure described by Hoffman (9). Dye dilutions of erythrocytes at 10⁶ cells/ml to 1/100 and extracts of Dr. Leescher diluted 1/10 to 1/1000 were placed at 4°C. The cultures were transferred to a 37°C water bath, incubated 30 minutes with stirring and sonicated at 5000g for 10 minutes at 4°C. The control cells were passed over them, and standardized to the desired concentration in the aqueous buffer.

3D inhibitor synthesis. 3D inhibitor synthesis was conducted using the GCPD subtractive assay described by Jeffcoat (199). These synthetases in the Zimmerman state (GCPD + NAD⁺) were mixed with an equal volume of 3D inhibitor solution diluted in 50 mM Tris. The reaction mixture was incubated at 20°C for 10 minutes with constant stirring.

After 10 min, three volumes of ice-cold complement diluted 1/25 in 0.1M KCl (0.02-0.05 ml) were added. The tubes were then incubated for 60 min more at 37°C with shaking. At the end of the incubation period, 10 μ l of ice-cold PBS was added to each reaction mixture. The cells were centrifuged at 5000 rpm for 10 minutes at 4°C and the cellular supernatant of the supernatant fluids were determined in a wave length of 411 nm.

Depletions. 10000 volumes of monomer ethyleneglycolic acetate (MGA, Fisher Scientific Co., Fair Lawn, NJ) and ethyleneglycolic (carboxymethyl ester) 1,2-hexamethylene diol (HED, Sigma Chemical Co., St. Louis, MO) were prepared as described by Fine et al. (11). 1% stock solutions were stored at 4°C and diluted to a final concentration of 200 μ l before use. Reversion (RV) was prepared as described by Fine et al. (11).

Complement consumption. The ability of erythrocytes treated with LPF and/or 24 U/ml Htr to consume complement was determined in reaction mixtures containing 0.1 ml of the treated cells (1.0 \times 10⁶ cells) in 1.0 ml and 0.1 ml of normal or absorbed guinea pig serum diluted with either 50% or 90%. The cultures were incubated with shaking 40 minutes at 37°C. Following the incubation period, the cells were pelleted at 5000 rpm for 10 minutes. The supernatant fluids were immunoprecipitated with digoxigenin labelled rabbit and were analyzed for membrane attack complex activity using a modification of the procedure as outlined by Salter and Royst (21).

Antimicrobial activity of LPS and its fragments. Repeated titrations of *Salmonella* typhimurium LPS at concentrations ranging from 0.001 μ g to 80.0 μ g/ml as measured with polyvinyl and bovine tracheobronchial antisera, employing erythrocytes at 2.0 \times 10⁶ cell/ml were carried out. The results of a representative experiment employing polyvinyl antisera are given in table III. It can be seen that among the LPS isomers the erythrocyte-coating capacity is a reversible action. Additionally, a number of μ g/ml bacterial cell wall mycobacterial were exposed to 0.001 to 0.20 μ g/ml of yeast LPS. Therefore, an optimal coating with 0.01 μ g/ml yeast LPS (defined as the incidence of the greatest efficiency of LPS enhancing complete hemolysis) by either polyvinyl or bovine antisera was found to 0.10 μ g/ml. Bovine erythrocyte bovine antisera to yeast LPS polymer have optimal coating doses of 0.10 μ g/ml and 1.00 μ g/ml, depending on the age of the antisera. These results are identical for bovine and sheep erythrocytes. LPS extracted by the Margalit procedure resulted in an optimal coating dose of 0.20 μ g/ml as was determined with bovine and sheep antisera.

LPS receptor activity, as evaluated in sheep plasma, was based on the ability of a given erythrocyte preparation to inhibit the complete lysis of an optimal coating with of LPS onto either sheep or bovine

Influence of various LPS concentrations
and NP Capping Rates (polymer)

LPS (μ g/ml)	Dose ^a	
	uncoated	coated ^b
0.0	80	80
0.05	110	80
0.1	100	80
0.2 ^c	40	80
0.5 ^c	10	100
1.0 ^c	10	100
2.0 ^c	10	100
5.0 ^c	10	80
0.15	10	80

The reciprocal of the dilution of anti-LPS serum affording maximal hemagglutination.

^a Stock solutions of LPS (1.0 mg/ml PBS) were diluted (10%) for these assays.

synthesizes. Table (11) which summarizes the results obtained with several crude human placental preparations of erythrocyte extracts. These data indicate that the range of LPS-receptor concentrations or dilutions needed to yield optimum inhibition of LPS erythrocyte casting varied with the source, concentration and conditions (salient) of the erythrocyte extracts extraction procedure.

(iii) *Inhibition activities of crude human extracts of lung and skin erythrocyte extracts*. DACTE measurement by a crude erythrocyte extract, prepared by the procedure described in the section on materials and methods is shown in Figure 5. This procedure was used to determine the inhibitory activity of such extracts. Color controls for the presence of hemoglobin in the higher concentrations of crude preparations were necessary. The reciprocal of the dilution of a crude extract preparation yielding greater than 50% inhibition of the lysis of DACTE by LPS are also shown in Table (11). These results clearly indicate that LPS-receptor and $\text{L}\alpha$ inhibitor activities are centered on what the authors term *crude human erythrocyte extracts* obtained by either Sykes and/or Bellman's procedures. Of interest also, is that the potency of the two activities varied to the same extent.

A comparison of the pharmacological properties of the $\text{L}\alpha$ inhibitor and LPS-receptor from erythrocyte extract extracts. The above data suggests that the LPS-receptor and $\text{L}\alpha$ inhibitor are either identical or closely related molecules, therefore, additional evidence to resolve this issue was sought. Human and sheep erythrocyte extracts were subjected to and fractions of Sykes and Bellman's procedures (16, 17) and

Table 1

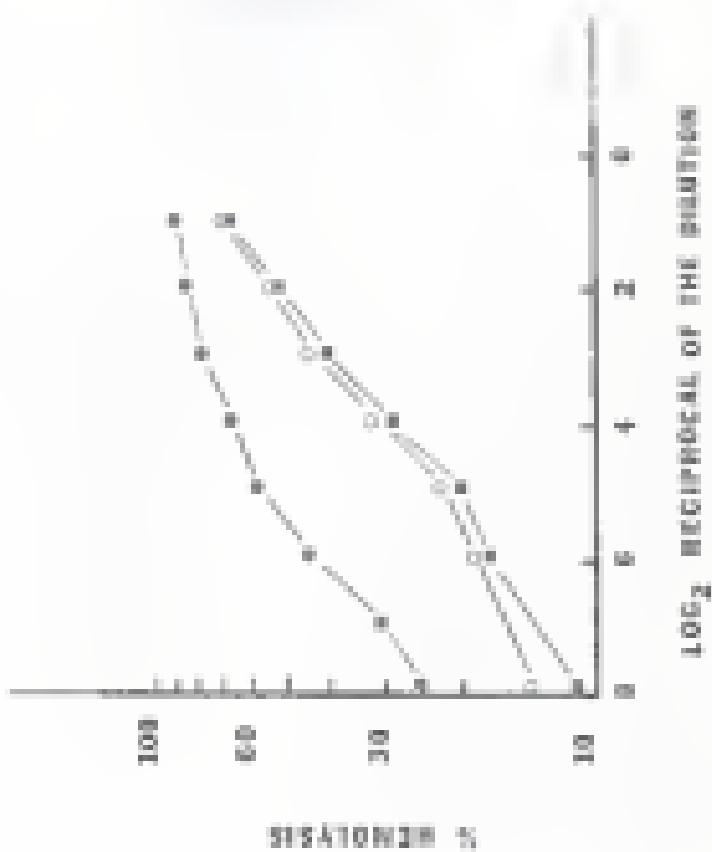
Comparison of L-Succinyl and D-Succinyl Derivatives of Tryptophane Derivatives

Drug Substance (Sigma)	Tryptophane Derivatives	
	IC_{50} (μM) ^a	IC_{50} (μM) ^b
None cells, untreated		
at pH 2.0 (Sarcoma)		
Drug resistant bacteria phase	40	40
Drug sensitive phase	200	200
High (2000) 1000 (Sar)	200	200
High (2000) 1000 (Sar)	40	40
None cells, infected		
at pH 2.0 (Sarcoma)		
Drug resistant bacteria phase	200	100
None cells, projected		
at pH 2.0 (Sarcoma)		
Drug resistant bacteria phase	0	0
Drug sensitive phase	0	0
Drug sensitive phase	0	0
Drug sensitive phase	0	0

^aThe methanol extract (100μg) shows complete inhibition of LPS activity.

^bThe inhibition of the granule effusion assay (IC₅₀) measured of DCTP (100 μM).

^cWith yeast extracts were measured by the proliferation of the drug resistant bacteria phase, 100 μM for 2 hours.



In Figure 2, Figure 4, the 1111 fractionation profile on Sephadex G-25, of the crude high speed top layer obtained from the crude rat liver methionyl-creatine preparation. Fractions were monitored at 280 and 310 nm and were assayed for LPS-receptor and 21 inhibitor activities as previously described. The peaks were observed with both activities eluting in the peak following the first elution. Close examination of the separation of profile indicates that there is a slight anaphase of the 21 inhibitor activity to the left of the LPS-receptor activity. This would suggest that perhaps the two activities may be different.

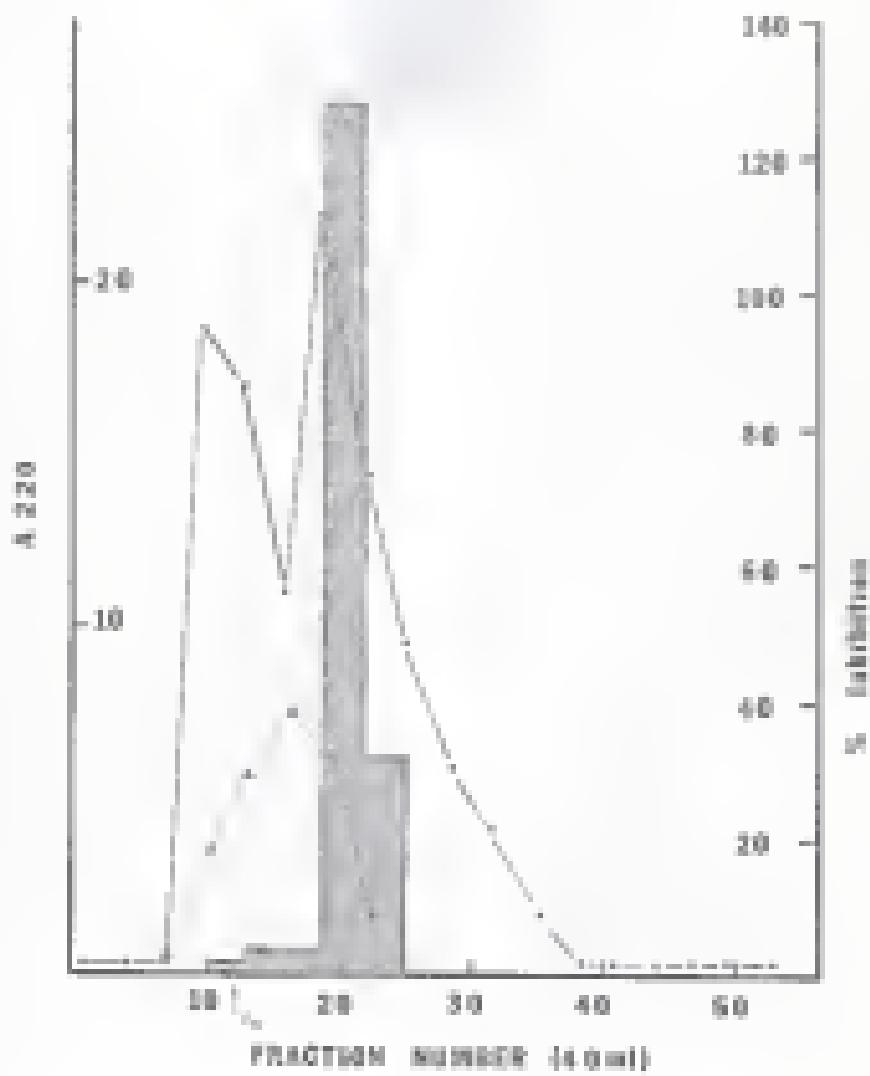
Further attempts to separate and purify the two activities were discontinued using the available chromatography. The aqueous 21 inhibitor peaks were pooled, dialyzed against the starting Tris-HCl buffer, and applied to a DEAE-Sephadex column. Fractionation was accomplished with a linear NaCl gradient. A radial chromatogram of the serial 10% pooled material [11] is shown in Figure 3. LPS-receptor and 21 inhibitor activities eluted in a relatively strong peak at about 0.3 M NaCl, again with the 21 inhibitor slightly preceding the LPS-receptor activity.

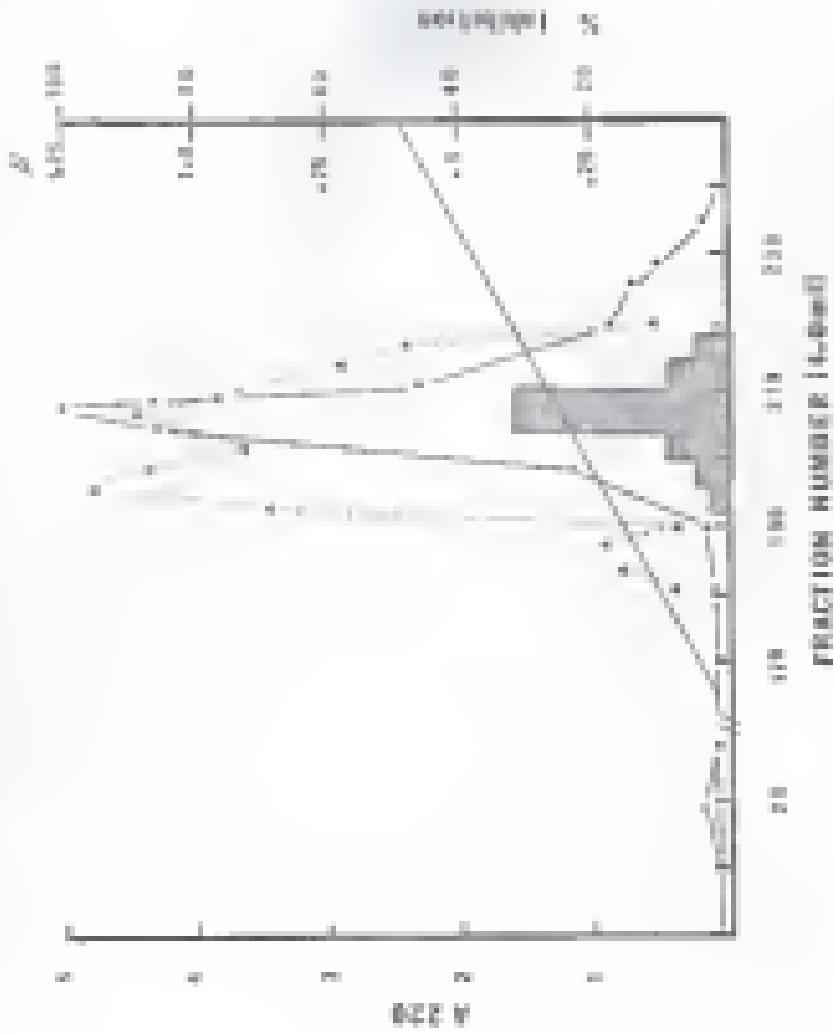
The recovery of the LPS-receptor and 21 inhibitor activities following aqueous 21 and DEAE-Sephadex chromatography is presented in Table 17. It should be noted that gel filtration on Sephadex G-25 yielded only about 50% loss in the purity of both activities with recovery of only 65% of the LPS-receptor activity and 40% of the 21 inhibitor activity. DEAE-Sephadex was shown to result in a much as a 10-fold increase in activity, but resulted in a recovery of 80% of the specific LPS-receptor activity but only recovered 40% of the

Dose (mg/kg)	Mean (SD) % of control (n = 6)		P-value
	100	200	
Control	100 (0)	100 (0)	0.99
100	100 (0)	100 (0)	0.99
200	100 (0)	100 (0)	0.99
400	100 (0)	100 (0)	0.99
800	100 (0)	100 (0)	0.99
1600	100 (0)	100 (0)	0.99
3200	100 (0)	100 (0)	0.99
6400	100 (0)	100 (0)	0.99
12800	100 (0)	100 (0)	0.99
25600	100 (0)	100 (0)	0.99
51200	100 (0)	100 (0)	0.99
102400	100 (0)	100 (0)	0.99
204800	100 (0)	100 (0)	0.99
409600	100 (0)	100 (0)	0.99
819200	100 (0)	100 (0)	0.99
1638400	100 (0)	100 (0)	0.99
3276800	100 (0)	100 (0)	0.99
6553600	100 (0)	100 (0)	0.99
13107200	100 (0)	100 (0)	0.99
26214400	100 (0)	100 (0)	0.99
52428800	100 (0)	100 (0)	0.99
104857600	100 (0)	100 (0)	0.99
209715200	100 (0)	100 (0)	0.99
419430400	100 (0)	100 (0)	0.99
838860800	100 (0)	100 (0)	0.99
1677721600	100 (0)	100 (0)	0.99
3355443200	100 (0)	100 (0)	0.99
6710886400	100 (0)	100 (0)	0.99
13421772800	100 (0)	100 (0)	0.99
26843545600	100 (0)	100 (0)	0.99
53687091200	100 (0)	100 (0)	0.99
107374182400	100 (0)	100 (0)	0.99
214748364800	100 (0)	100 (0)	0.99
429496729600	100 (0)	100 (0)	0.99
858993459200	100 (0)	100 (0)	0.99
1717986918400	100 (0)	100 (0)	0.99
3435973836800	100 (0)	100 (0)	0.99
6871947673600	100 (0)	100 (0)	0.99
13743895347200	100 (0)	100 (0)	0.99
27487785694400	100 (0)	100 (0)	0.99
54975571388800	100 (0)	100 (0)	0.99
109951142777600	100 (0)	100 (0)	0.99
219902285555200	100 (0)	100 (0)	0.99
439804571110400	100 (0)	100 (0)	0.99
879609142220800	100 (0)	100 (0)	0.99
1759218284441600	100 (0)	100 (0)	0.99
3518436568883200	100 (0)	100 (0)	0.99
7036873137766400	100 (0)	100 (0)	0.99
14073746275532800	100 (0)	100 (0)	0.99
28147492551065600	100 (0)	100 (0)	0.99
56294985102131200	100 (0)	100 (0)	0.99
11258997020426400	100 (0)	100 (0)	0.99
22517994040852800	100 (0)	100 (0)	0.99
45035988081705600	100 (0)	100 (0)	0.99
90071976163411200	100 (0)	100 (0)	0.99
18014395232682400	100 (0)	100 (0)	0.99
36028790465364800	100 (0)	100 (0)	0.99
72057580930729600	100 (0)	100 (0)	0.99
14411516186145600	100 (0)	100 (0)	0.99
28823032372291200	100 (0)	100 (0)	0.99
57646064744582400	100 (0)	100 (0)	0.99
115292129489164800	100 (0)	100 (0)	0.99
230584258978329600	100 (0)	100 (0)	0.99
461168517956659200	100 (0)	100 (0)	0.99
922337035913318400	100 (0)	100 (0)	0.99
1844674071826636800	100 (0)	100 (0)	0.99
3689348143653273600	100 (0)	100 (0)	0.99
7378696287306547200	100 (0)	100 (0)	0.99
14757392574613094400	100 (0)	100 (0)	0.99
29514785149226188800	100 (0)	100 (0)	0.99
59029570298452377600	100 (0)	100 (0)	0.99
118059140596904755200	100 (0)	100 (0)	0.99
236118281193809510400	100 (0)	100 (0)	0.99
472236562387619020800	100 (0)	100 (0)	0.99
944473124775238041600	100 (0)	100 (0)	0.99
1888946249550476083200	100 (0)	100 (0)	0.99
3777892499100952166400	100 (0)	100 (0)	0.99
7555784998201904332800	100 (0)	100 (0)	0.99
1511156999640380866400	100 (0)	100 (0)	0.99
3022313999280761732800	100 (0)	100 (0)	0.99
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245199238189092230047304658521600723728000	100 (0)	100 (0)	0.99
4903984763781844600946093170432001447568000	100 (0)	100 (0)	0.99
9807969527563689201892186340864002895136000	100 (0)</		

What's more, according to the new study, the more people you have in your life, the more you're likely to be healthy. The researchers found that people with more social connections were less likely to die from heart disease, stroke, or cancer.

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activities of tritiated activity. A comparison of the α -filtration and DAB-deprotein extract profiles (Figures 4 and 5) and the data in Table III, revealing a 100-fold in the total area but with a 16-fold increase in activity, indicate good purification of the α -activities. These data do, however, suggest further differences in the two activities by the differences in their respective behaviors following treatment III and DAB-deprotein chromatography.

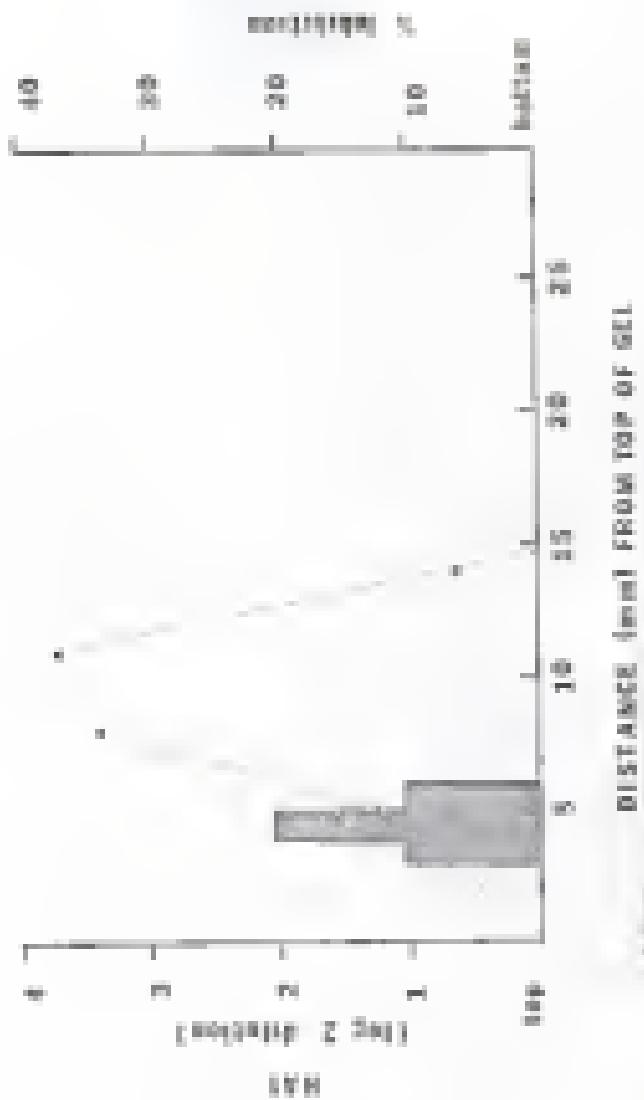
The transparency of the active DAB-deprotein fractions was checked by electrophoresis on 7.5% polyacrylamide gels at pH 8.0. As shown in Figure 6, 4–12 protein bands were observed on the crude extracts with a sharp band and a diffuse band appearing just behind it at the top of the gel when partially purified DAB-deprotein fractions were electrophoresed. Analysis of diphoresis gels indicated that most of the LP-enzyme activity was localized in an area about 6 mm below the gel with the 10-kilodalton activities spread over a forty-mm area at the top of the gel with a peak of activity of about 11.0 mm. (Figure 6)

10. Tritiated-gel LP-enzyme activity of sheep erythrocyte extracts. The reverse phase of bovine erythrocyte sheep erythrocyte extracts, prepared by Springer's extraction procedure at pH 8.0 and at pH 9.0, but neither deprotein LP-enzyme nor LP-enzyme activities (LP-enzyme activity between 10 inhibitor units, however, observed in an other soluble fraction of the solution) could be observed in Table III.

Figure 1.

Correlation between the distribution of HPMs and
the number of HPMs per unit area

Figure 1 shows the correlation between the distribution of HPMs and the number of HPMs per unit area. The figure consists of two panels. The left panel is a scatter plot showing the relationship between the number of HPMs per unit area (Y-axis, 0 to 100) and the distribution of HPMs (X-axis, 0 to 100). The data points show a strong positive correlation, with a regression line fitted to the data. The right panel is a histogram showing the distribution of HPMs per unit area. The X-axis represents the number of HPMs per unit area, ranging from 0 to 100. The Y-axis represents the frequency, ranging from 0 to 10. The histogram shows a unimodal distribution with a peak around 50 HPMs per unit area.



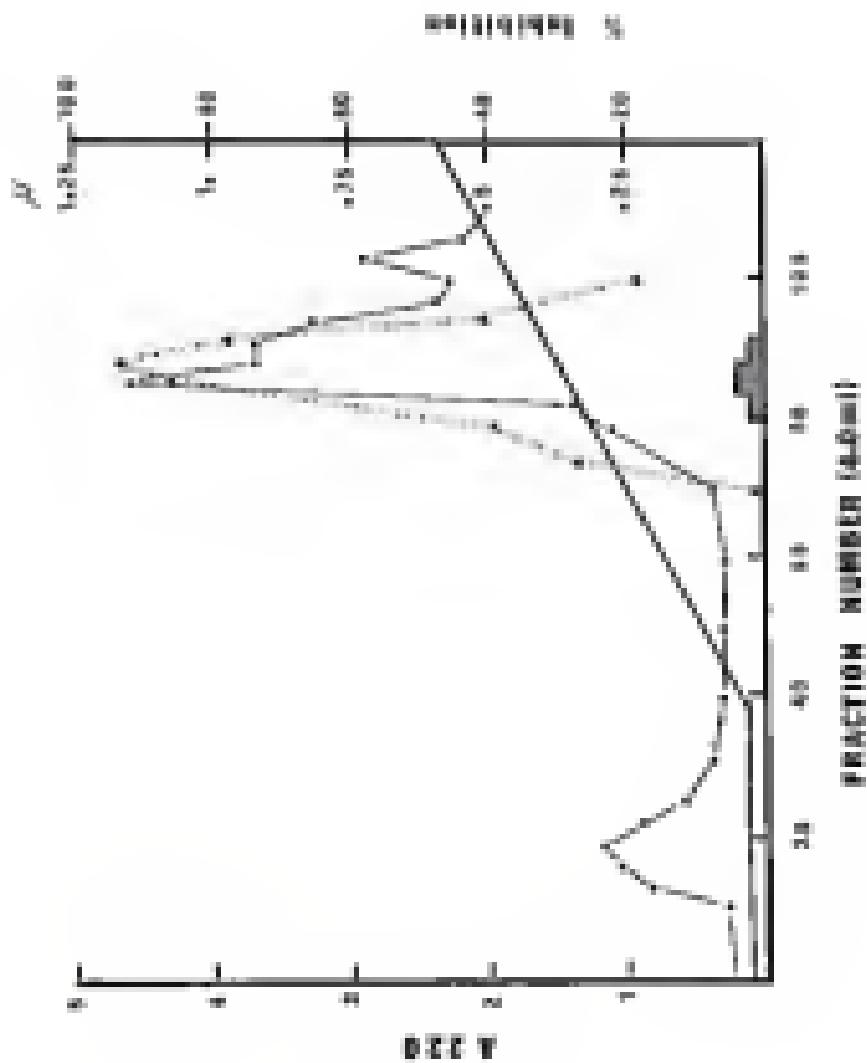
A second class of difference in the two extractions was evident when it was previously observed that a shift in the pH from 8.0 to 6.0 in the buffer extraction step of the crude erythrocyte lysates resulted in a progressive quenching of UTH receptor activity in detectable affinity, but little loss in receptor activity. The elution profile of the UTH-receptor activity following the exchange chromatography on DEAE columns of the partially purified material, as shown in Figure 7, was similar to that observed for material extracted at pH 8.0. UTH-receptor activity assays were not carried out on the recovered buffer eluates because these materials had been discarded before the onset of these observations were realized.

A shift in the pH to 6.0 of a crude tetanol extract of erythrocyte lysates obtained at a pH of 8.0 affected neither the UTH receptor nor the receptor activities. This suggested the possibility that the UTH-receptor activity of the material extracted at a pH of 8.0 was not destroyed but was probably transferred into another phase. UTH receptor assays carried out at pH 6.0 employing buffer eluates of erythrocyte lysates, using borate's presence, in an attempt to baseline the UTH-receptor activity. As shown in Table 8, some UTH receptor activity was observed in the crude lysate tetanol phase when 50% of packed columns were eluted. There was no detectable UTH receptor in this tetanol layer. The lipid phase was eluted with 10 volumes of H2O and further suggested to be either extraction resulting in three layers a pellet, an aqueous layer, lipid interface, and an organic layer. 50% of the UTH-receptor activity was recovered in the organic ether layer.

Favorable		Unfavorable		Total	
Response	Percentage	Response	Percentage	Response	Percentage
Very favorable	10.0	Very unfavorable	10.0	Very	10.0
Somewhat favorable	30.0	Somewhat unfavorable	30.0	Somewhat	30.0
Unfavorable	50.0	Unfavorable	50.0	Unfavorable	50.0
Total		Total		Total	
Favorable		Unfavorable		Total	
Very favorable		Very unfavorable		Very	
Somewhat favorable		Somewhat unfavorable		Somewhat	
Unfavorable		Unfavorable		Unfavorable	

mentally, a significant shift occurred to a more positive attitude towards the use of alcohol in the United States.

1971-1972. See, *Journal of Latin American Studies* 10, 1978, pp. 484-511. Another view is that, though *lucha* may be a more appropriate term than *revolution* in this context, the term *revolution* is not inappropriate in the sense that it is used in the literature on Latin American revolutions. The term *revolution* is used here to denote a process of social and political transformation that, while it may not be a revolution in the sense of a violent overthrow of an existing political order, nevertheless creates a new political order. The term *revolution* is used here to denote a process of social and political transformation that, while it may not be a revolution in the sense of a violent overthrow of an existing political order, nevertheless creates a new political order.



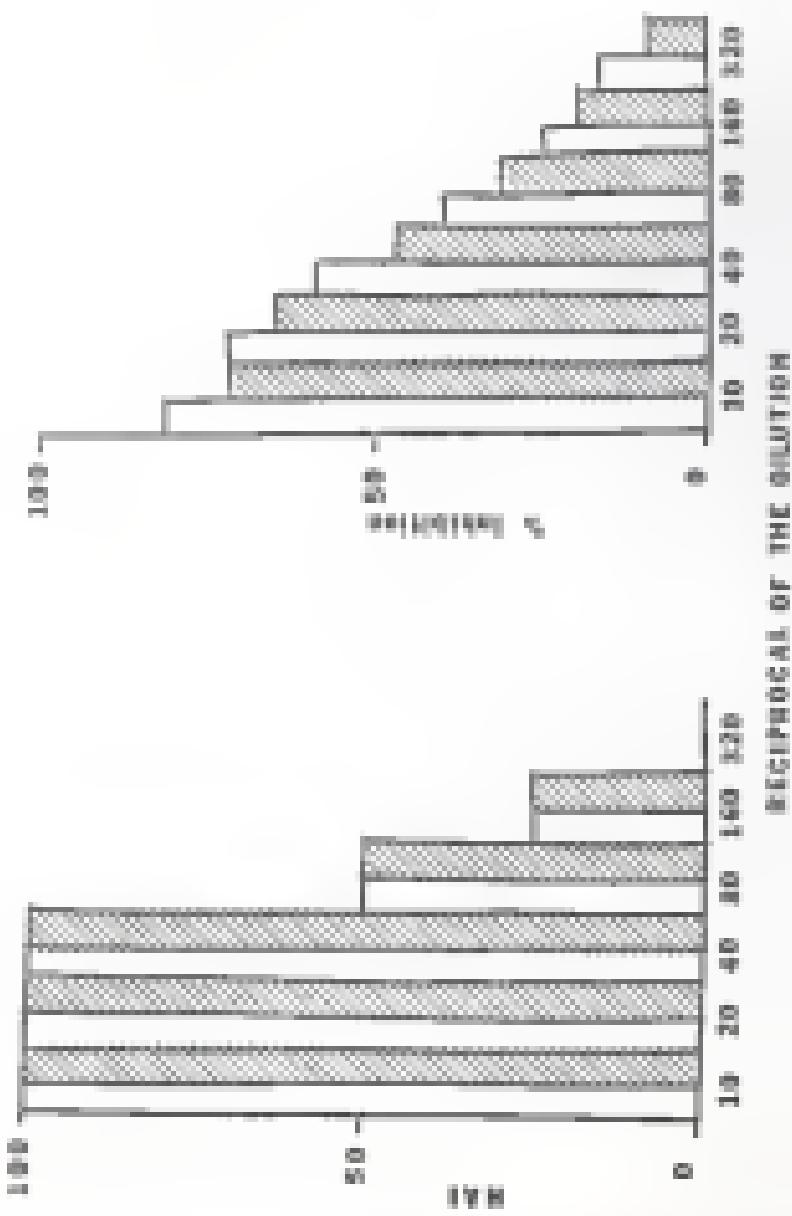
10³), the β -D-glucuronidase activity (10000 U/mg), and β -D-glucuronidase. A comparison of the peritrophic bodies (P and P₀) and their extracts obtained at pH 6.0 and pH 5.0 indicates that, as was observed at pH 6.0, extracts obtained at pH 5.0 and subjected to SDS-PAGE chromatography resulted in a substantially greater loss of total hemolysin activity as estimated by the absorbance at 280 nm and the yield of β -D-glucuronidase activity.

Treatment of *Brucella*'s crude lysed extracts with shear conditions. To further establish that the two activities are differentially affected, equal volumes of shear erythrocytes at either 10³ or 10⁴ cell/ml were mixed with equal volumes of a lysed fresh sand bee (Lys) extract prepared according to the procedure of Berlese. This extract has an initial LPS-receptor titer of 128 and a β -D-glucuronidase titer greater than 20. Control tubes consisting of equal volumes of buffer and the extracts were also prepared. All tubes were mixed at 30°C for 30 minutes with shaking. The cells were pelleted by centrifugation and the supernatant fluids along with the buffer controls were eluted and assayed for LPS-receptor and β -D-glucuronidase activities. As can be seen in Figure 8, the β -D-glucuronidase activity was reduced substantially when extracts were treated with 10³ cell/ml. In contrast, the LPS-receptor activity remained constant when the cell treated and buffer control supernatants fluids were compared.

The biological consequences of the β -D-glucuronidase and LPS-receptor in the *Brucella* membrane. The data in the previous section indicated two associated events. First, the membrane of the human erythrocyte,

Figure 4

Treatment of normal epithelial monolayer in monolayer culture system prepared by the method of Springer. Either 10⁶ these epithelial cells/ml or 10⁵ fibroblasts/ml were mixed with an equal volume of the untreated culture. Following a 24 hour incubation period at 37°C, the cell suspension was pelleted and cell supernatant fluids were assayed for U6-uridylate and D-ribofuranose activities. The hatched bars represent the monolayer system treated solution, and the open bars the buffer-treated controls.



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Die beiden ersten Verszeilen sind in der handschriftlichen Fassung von 1801 (vgl. oben) nicht erhalten, während die dritten beiden Verszeilen in der handschriftlichen Fassung von 1801 (vgl. oben) nicht erhalten sind.

19. *Leucosia* (Leucosia) *leucostoma* (Fabricius) (Fabricius, 1775: 116. Type locality: India).

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Group	Mean	SD	Median	Range
Control	1.00	0.00	1.00	0.00-1.00
100 mg/day	1.00	0.00	1.00	0.00-1.00
200 mg/day	1.00	0.00	1.00	0.00-1.00
400 mg/day	1.00	0.00	1.00	0.00-1.00
800 mg/day	1.00	0.00	1.00	0.00-1.00
1600 mg/day	1.00	0.00	1.00	0.00-1.00
3200 mg/day	1.00	0.00	1.00	0.00-1.00
6400 mg/day	1.00	0.00	1.00	0.00-1.00
12800 mg/day	1.00	0.00	1.00	0.00-1.00
25600 mg/day	1.00	0.00	1.00	0.00-1.00
51200 mg/day	1.00	0.00	1.00	0.00-1.00
102400 mg/day	1.00	0.00	1.00	0.00-1.00
204800 mg/day	1.00	0.00	1.00	0.00-1.00
409600 mg/day	1.00	0.00	1.00	0.00-1.00
819200 mg/day	1.00	0.00	1.00	0.00-1.00
1638400 mg/day	1.00	0.00	1.00	0.00-1.00
3276800 mg/day	1.00	0.00	1.00	0.00-1.00
6553600 mg/day	1.00	0.00	1.00	0.00-1.00
13107200 mg/day	1.00	0.00	1.00	0.00-1.00
26214400 mg/day	1.00	0.00	1.00	0.00-1.00
52428800 mg/day	1.00	0.00	1.00	0.00-1.00
104857600 mg/day	1.00	0.00	1.00	0.00-1.00
209715200 mg/day	1.00	0.00	1.00	0.00-1.00
419430400 mg/day	1.00	0.00	1.00	0.00-1.00
838860800 mg/day	1.00	0.00	1.00	0.00-1.00
1677721600 mg/day	1.00	0.00	1.00	0.00-1.00
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26843545600 mg/day	1.00	0.00	1.00	0.00-1.00
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1826875450282297071404943829251				

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Spurzahnkäfer (2000) in der Natur, von J. H. J. B. Schipperijn et al. (2001) und in der Wildnis (2001) von J. H. J. B. Schipperijn et al. (2001) in Europa. Es ist zu hoffen, dass die Ergebnisse der vorliegenden Arbeit die Ergebnisse der anderen Studien bestätigen werden.

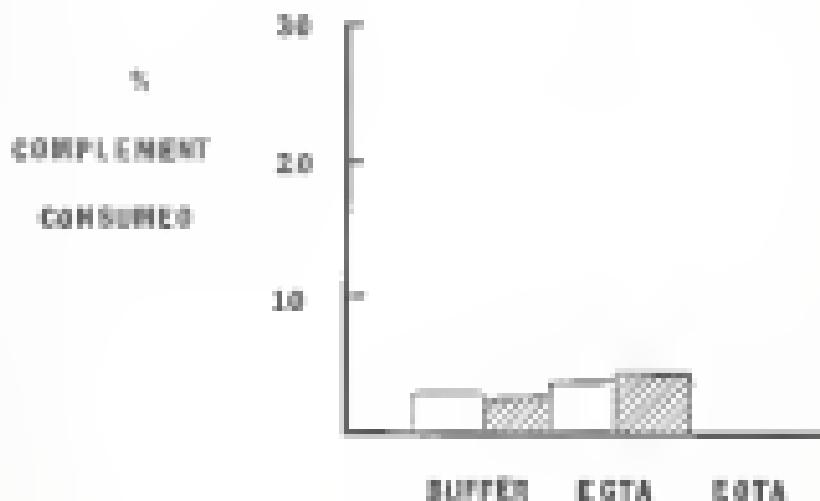
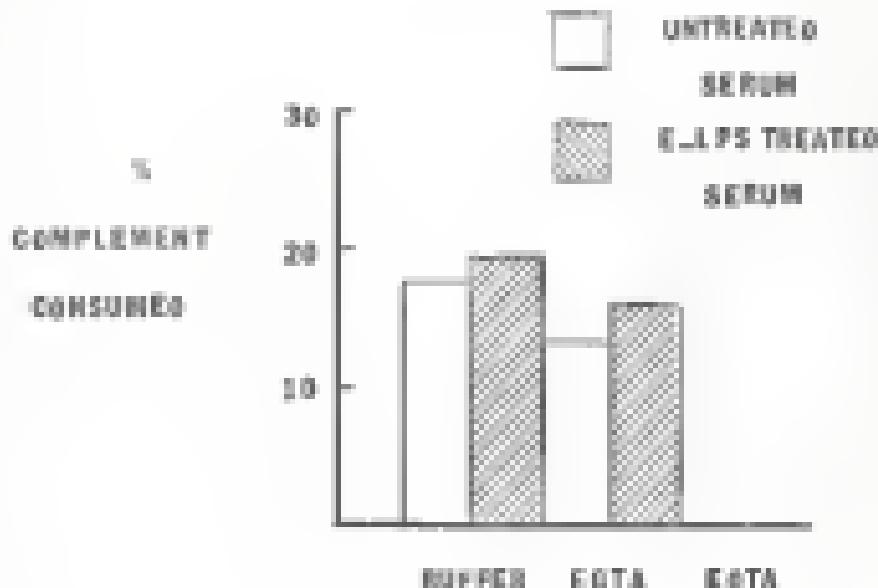
which (i) mainly reacted to coagulate mediated Lytic pathways, leaving the apparently different molecules with biologically exerting properties: IgG1-particles with a high affinity for C3b-
Bb-convertase which are potent activators of the complement system, and a class of molecules shown to be potent inactivators of complement. Second, sheep cells which are normally extremely sensitive to human Lytic素 have been shown to be devoid of the Bb-activator for human molecules with an affinity for C3 which are confined to the IgG1-
activity of the molecule.

As shown in Figure 3, the interaction of free C3b-biotinylated by both the Biotin and Biotinyl-proteinase with guinea pig serum which had been absorbed with sheep IgG reacted with C3b (30.4%), resulted in a substantial consumption of complement. Additionally, it can be seen that C3b (Biotin) appeared to be a much more efficient activator of the alternative pathway pathway compared to C3b activated by the Biotinyl-proteinase.

C3b (activated by both mechanisms) reacted with the surfaces of sheep erythrocytes showed a similar profile (Table III), except erythrocytes reacted with C3b activated by the Biotinyl-proteinase were the most efficient activators of complement in the absence of natural activators to C3b. It is of interest, therefore, to determine if C3b or the surface of sheep erythrocytes, in the presence of the Bb-activator and Bb-C3b-activated guinea pig serum, could alter the alternative consumption profile of C3b-C3b to explore this possibility, erythrocytes were reacted with C3b and Bb-activator that reacted with guinea pig serum (aspirated

TABLE 4
**Comparison of Total Complexity (Jillings) in D-IPMS
 or D-IPMS/absorbent
 for Extracted and IPMS-Carried Sheep Erythrocytes**

Cell Suspension	Average Recovery	
	D-absorbed series	D-IPMS-absorbed series
1	—	—
D-IPMS-carried	99.1	10.08
D-IPMS-free	42.4	6.38

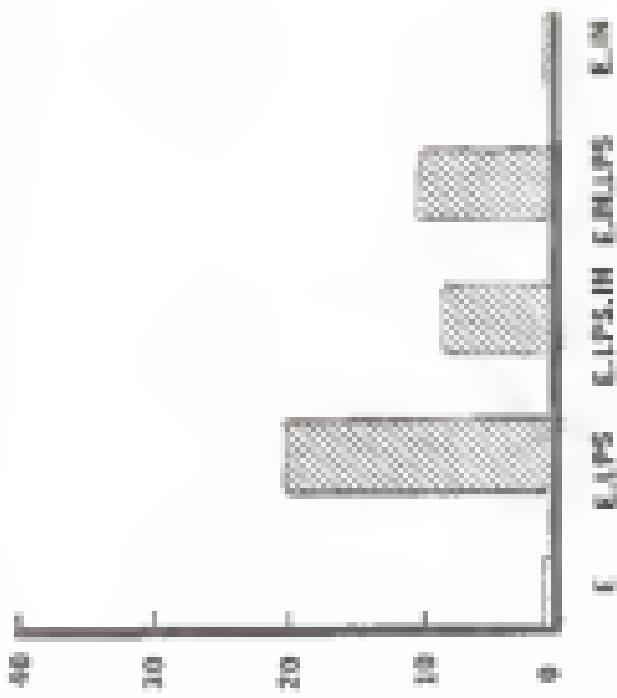


After 24 h, 10,000 cells/well of cells were prepared. One group was coated with LPS only. In addition, a second group was coated with LPS first, then was treated with the IgG-inhibitor (0.45% IgG), a third group was treated with IgG-inhibitor first, followed by the LPS-receptor (G + IgG), a fourth group was treated with IgG-inhibitor only (0.45%), and a fifth group of cells (H) was treated with PBS under the same conditions and served as the control. The efficiency of the LPS-coating procedure in the presence and absence of the IgG-inhibitor was evaluated by employing the five groups of cells and their respective supernatant fluids for LPS activity, employing the hemagglutination assay (described in Bernath and Petrusch) as determined with antisera to *Salmonella typhimurium* strain B. The results of these assays indicated that all of the LPS-treated cells adsorbed equal quantities of LPS to the presence and absence of the IgG-inhibitor. The confluent-concentration profiles of the five groups of erythrocytes treated with 0.45% (w/v) dialyzed gelatin 10 μ g serum are presented in Figure 10. It can be seen that the presence of IgG-inhibitor on the cell surface does not contribute to the LPS-mediated hemolytic action of complement. Of particular interest was the fate of the cells when confluent were treated with IgG + LPS-coated erythrocytes, significantly less cells were lysed. This was in contrast to the case where confluent were coated with 0.45% and the cells were completely lysed. These results suggested the overall IgG-inhibitor complement activation had been intact, but that cells were protected from lysis by the presence of IgG-inhibitor on the material.

Figure 1b

Preparation of complexes in U-101 treated fibrotic patients ($n=10$) versus by U-101 under U-101 free control fibroblast-myoblasts. The death of constituting 10^6 fibroblast-myoblasts (U-101) either untreated or treated with U-101 (U-101), U-101+U-101 or U-101 and U-101 free (U-101-U-101) were measured when 0.1 ml of culture was carried at 37°C. For 1 hr fibrotic complexes myoblasts activities were assayed and the % of the available complement contained was calculated.

COMPLEXITY
COHESIVENESS



DISCUSSION

The experiments reported here have demonstrated that relatively free human erythrocyte membranes possessing LPS-receptor activity described by the method of Springer et al. (16) were also capable of initiating complement mediated lysis. The anticomplementary activity of these extracts was demonstrated to share many of the properties of the LPS inhibitor previously described by Hartman (10).

Data are presented which strongly suggest that the two biological activities are closely associated, but separate. Evidence for this has provided by the results of the different experimental approaches to the problem of Springer human erythrocyte stromal extracts. The first has been to the chromatographic separation of reference LPS and LPS-deglycanated where slight differences between the elution patterns and recoveries of the two activities were observed. The second piece of evidence came from the electrophoretic profile of the crude and partially purified extracts on 2.5% polyacrylamide gels under non-denaturing conditions. The LPS inhibitor activity was shown to cover a fairly large area at the top third of the gel with the LPS-receptor activity being localized in a narrow, single band with a peak of activity near the top of the gel. A third line of argument was based on the modification and separation of the two activities into different phases with the pH during the crude stromal lysed erythrocyte procedure

varies from 0.2 to 0.3. The fourth approach, based on the high affinity of one β -galactosidase for the membranes of sheep erythrocytes, demonstrated that the D-galactosidase activity was partially removed leaving the LPS-receptor activity unchanged by the treatment of the intestinal extracts with sheep erythrocytes. Finally, evidence was presented indicating that the binding specificity of sheep erythrocytes for LPS of *gram-negative* bacteria is limited to a large majority of the crude bacterial extracts and to none of all detectable D-galactosidase activity.

It should be emphasized, however, that these experiments cannot exclude the possibility that both activities may be associated on the same macromolecules with the differences reported here being a consequence of experimental manipulations. That the two activities may be a function of a single macromolecule is certainly a more possibility. Springer et al. (19) in assessing the chemical and physical properties of a homopurified preparation of the LPS-receptor, observed that both ethylmagnesium and dissociating polyacrylamide gel electrophoresis under standard conditions yielded two fragments, one of which displayed significantly only at 310 nm. Reconstitution of the ethylmagnesium fragment restored high LPS-receptor activity to only one of the fragments. These studies are very suggestive and do not permit a decision as whether the activities are on the same macromolecule.

In contrast, the data obtained from sheep erythrocytes which completely lack D-galactosidase, but which possess LPS-receptor activity, would support the theory that the two macromolecules may be distinctly different. However, the evidence would support that the LPS-receptor

on these cells differ from those observed on bone cells, since they are affected by the T-cell subset.

Good generalities of the LPH-receptor and D1 inhibitor activities following GM-CSF-dependent differentiation is reflected by the quantitative data tables. These results would suggest, however, that a pre-judicial purification step of should be modified to enhance the purity of the two activities.

The observation that these erythrocytes possess molecules with receptor specificity for LPS is not surprising. For D1 having been shown that these cells could be sensitized by the presence of LPS of *gram-negative* bacteria and that these sensitized cells D1 readily bind to the presence of antibodies antiserous to LPS and complement (40). In contrast, however, it was not observed when LPS treated bone erythrocytes were treated under the same conditions. This refutes the possibility that bone erythrocytes, *in vitro*, lack of the D1 T-lymphocyte when treated with LPS, are extremely resistant to LPS mediated lysis because of the presence of inhibitor molecules.

The findings reported here generally agree with those of Rittling et al. (41) indicating that LPS treated bone erythrocytes can activate the complement system in the presence of certain antibodies to the LPS. In contrast to their results, however, erythrocytes coated with a preparation of LPS adsorbed by the procedure of Kortestol were shown to be capable of activating the complement system in the absence of natural antibodies to LPS. Additionally, that this LPS adsorbed by the Kortestol (LPS-adsorber) was shown to be a more effective adsorber of the

complement system than LPS extracted by the Bacterial procedure (LPS-antigenic) in the presence and absence of natural antibodies to LPS. However, since LPS extracted by the Davis becomes cell associated, its capacity to activate the complement system in the absence of natural antibodies is greatly diminished. This is significant because LPS activation of the complement system by an antibody independent mechanism requires either an exposed Mannose moiety or polysaccharide core (10,11). This would suggest then that the antigenicity of LPS due to the core may be different from that of (LPS-antigenic), resulting in the masking of active sites necessary for the activation of complement.

The fact that different preparations of LPS from the same organism when coated onto the surface of glass microtiter, activated the complement system to different degrees and for different pathways, depending on the presence or absence of natural antibodies to LPS, introduces the possibility that the LPS activation of complement may involve interactions other than the LPS molecule alone. This especially appears to be true when LPS extracted by the bacterial procedure, which was shown to activate complement in the presence and absence of natural antibody to LPS, is found to contain less protein and lipoproteins than LPS extracted by the Davis procedure.

Having the IgG inhibitor on LPS treated sheep erythrocytes reduced the ability of erythrocytes to consume complement. The fact that the cell Ig is protected even when complement is activated suggests that the Ig inhibitor may occupy specific areas on the red cell membrane rendering it resistant to protease lysis, but also having Ig antibodies

de-potentially inhibited complement lysis. This may be due either to the masking of LPS-receptors resulting in low LPS uptake (assumed to be to the one tenth of the presented LPS material on the cell membrane thus blocking the activation of the membrane attack system of complement). An analogous role in the human erythrocytes is already assumed by the D8 inhibitor resulting from cell naturally having no LPS mediated lysis.

The findings presented here have led us to hypothesize that a necessary criterion for resistance to LPS induced complement mediated lysis would be the localization of LPS-receptor and D8 inhibitor molecules on the same membrane. Thus would imply that any cell with devoid of D8 inhibitor molecules would be far more susceptible to the cytolytic action of LPS activated complement.

LPS of gram-negative bacteria are potent activators of the complement system. In the past clinical relevant, therefore, bacterial substances on the surface of erythrocytes which bind LPS are found closely associated with membrane capable of activating complement mediated lysis. The symptoms of several infectious diseases, such as typhoid fever, have been observed to include very intensive erythrophagocytic activity by macrophages of lymph nodes. The consequence of this observation could have great clinical importance. Erythrocytes coated with LPS, to interact with complement, would naturally tend to activation of the complement system followed by increased phagocytosis with infimal cytotoxicity. This would lead essentially to an amplification of the activities of the complement cascade resulting in either clearance or a heightened inflammatory response.

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After I left the University of Florida, I moved to Jacksonville, Florida. I met Shirley Anne Jackson, my roommate, who was attending Florida Institute, and the second of three children. Shirley and I attended and we graduated May 26, 1968, valedictorian of my class at Baylen High, a private school for girls, Jacksonville, Florida, Jacksonville, Florida.

I later attended Florida College, majoring in premedicine with a minor in psychology, graduating June of 1970. Lack of funds prevented Shirley and me from going to college, therefore, I was employed as a research technician at University of Florida for one year. I later earned the University of Kansas at Lawrence, Kansas to receive a Master's degree in Biochemistry specializing in molecular physiology. After a year and a half I completed the credit hour requirements for the master's degree and was employed as a research technician in the Department of Biochemistry, where I served as an assistant to a Research Scientist. In October 1973 I was married to Irvin Lee Jackson. Following which we relocated to Chicago, Illinois.

During my three years in Chicago, I worked as a research technician at the American Medical Association Biomedical Research Institute and later as assistant supervisor of the Biophysiology Research Department at the International Survey Institute. On May 16, 1976 I gave birth to a son, Robert O'Neill, following which we relocated to Parsippany, New Jersey.

In 1999, I moved to New Jersey. I was employed at the pharmaceutical company, Novartis where I worked in studies on the microbiology of *Chlamydia*. After relocating to Florida, I accepted and finally joined, Biogenics where my husband was employed as the president of the writing company.

Following a legal separation, I returned with my son, Joseph, to Novartis to serve a sabbatical at the University of Florida, majoring in Microbiology and specializing in Immunology. I am presently employed at Abbott Labs of North Chicago as a project manager in research and development. In addition to immunology, I enjoy tennis, chess, bridge, writing, and running.

I certify that I have read this study and that in my opinion it conforms to acceptable standards of scholarly presentation and is worthy, in form and quality, as a dissertation for the degree of Doctor of Philosophy


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